

GCCAGGTCTGGCACCATGCACTAGGATACCCAGAACGCTGCAAGGCCACGCC
 CTCCTCACTTCAGGGGTCCTCTCCCCATTGCCACCACCCACCATGGCTGGG
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 GTGGCGGGCCTGGGGGACCCTTGGTCATCGACAGCCGGTCCTTCGTGGAGTAT
 AACAGCTGCCACGTGCTGAGCTCTGTGAATATCTGCTGTTCAAAGCTGGTGAA
 GCGGCGCCTTCAGCAGGGAAAAGTGACAATTGCTGAGCTTATCCAGCCTGCTA
 CACGGAGCCAGGTGGATGCCACAGAACCACAGGATGTAGTGGTGTATGACCA
 GAGCACACGAGATGCCAGCGTGCTGGCAGCAGACAGCTTCCTGTCCATCCTGC
 TCAGCAAGCTGGACGGCTGCTTCGACAGTGTGGCCATCCTCACAGGAGGCTTC
 GCCACCTTCTCCTCCTGCTTCCCTGGCCTCTGTGAGGGCAAGCCTGCCACTCTA
 CCGTCCATGAGCCTCTCTCAGCCCTGCCTGCCTGTGCCCAGTGTTGGCCTGACC
 CGAATCCTGCCTCACCTCTACCTGGGCTCTCAGAAAAGATGTCTTGAACAAGGA
 TCTGATGACCCAAAACGGAATAAGCTATGTCCTCAATGCCAGCAACTCCTGCC
 CTA AACCGGACTTCATCTGTGAGAGCCGTTTCATGCGTATCCCATCAATGAC
 AACTACTGTGAAAAGCTGCTGCCCTGGCTGGACAAGTCCATCGAGTTTATTGA
 TAAAGCCAAGCTGTCCAGCTGCCAAGTCATTGTTCACTGTCTGGCTGGCATCTC
 TCGCTCTGCCACCATTGCCATCGCGTACATCATGAAAACCATGGGCATGTCTTC
 TGACGACGCATACAGGTTTGTGAAGGATCGGCGCCCCCTCCATCTCGCCCAACT
 TCAACTTCCTGGGCCAGTTGCTGGAGTATGAGAGGAGTCTGAAGCTGCTGGCT
 GCCCTGCAGACTGATGGACCTCACTTGGGGACCCCTGAGCCCCCTCATGGGCCC
 GGCAGCAGGCATCCCCTGCCCCGGCTGCCACCATCTACCTCAGAGAGCGCTG
 CCACTGGGAGCGAGGCAGCCACCGCAGCCAGGGAGGGCAGCCCAAGTGCTGG
 AGGGGATGCTCCGATCCCCAGCACAGCTCCAGCCACCAGCGCGCTGCAGCAG
 GGCCTGCGTGCCCTGCACCTCTCCTCTGACCGCCTCCAGGACACCAACCGCCT
 CAAGCGTTCCCTTTTCCCTGGACATCAAGTCGGCCTATGCACCCAGCAGGAGGC
 CCGACTTTCCCGGCCACCCGACCCCGGTGAAGCCCCGAAGCTCTGCAAGCTG
 GACAGCCCGTCTGGGGGCACACTGGGCCTGCCCTCGCCCAGCCCAGACAGCCC
 GGA CTCCGTTCCAGAGTGCCGCCCACGACCCCGCCGGCGACGCCCCCGGCTA
 GTTCGCCTGCCCGCTCCCCCGCGCATGGTCTGGGCCTGAACTTTGGAGACACG
 GCCCGGCAGACTCCACGGCACGGCCTCTCGGCCCTGTCGGCGCCCGGGCTGCC
 TGGCCCTGGCCAGCCGGCTGGCCCCGGGGGCTGGGTGCCGCCACTGGACTCCC
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 GGGTCCAGGCGCTGTGTTCTCCGCCTTTGGCCGGGTAAAGTGACGGCGCACCTG
 GACCCGGTAACAGCAGCAGCAGCGGTGGTGGTGGTGGTGGTGGTGGCGGCGG
 CGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGG
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 GGCAGCGTGAGGTCATCGAAGTATCGTGACCCTTCAGAAGTCCCTGTGCCCT
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 GGTA AATGGTTTTACTGCAATTTTTATCAAGAAGTAAATATTTTCGATTTTTTAT
 TTATTTAAGCTAGTGATCTGGCAACTGTGCGGGGCGGCCCTAAAGCTCTGTTTT
 TACTGTCTGGTATTTAAACTGAAACAGGTTTCTAAGCAATATGAGGCCACCTT
 CAATCCCAAACCTGGGTTGACAGGCCTGGGCCCCCTCCTTGCCCCCTCCCCTCTGG
 AAACATTACTGACCTTTCAAAGAGCTGCCAGCTTTCCTGCACTTTTTACATAA
 GAAAAAAGGGGGGGGGGGGAA (SEQ ID NO:1)

FIGURE 1

underlined = deleted in targeting construct

[] = sequence flanking Neo insert in targeting construct

[GCCAGGTCTGGCACCATGCACTAGGATACCCAGAACGCTGCAAGGCCACGCCCTCCTCAC
TTCAGGGGTCACTCTCCCCATTGCCCACCACCCACCATGGCTGGGGATCGGCTCCCGAG
GAAGGTGATGGACGCAAAGAAA] CTGGCCAGCCTGCTGCGTGGCGGGCCTGGGGGACCCTT
GGTCATCGACAGCCGTCCTTCGTGGAGTATAACAGCTGCCACGTGCTGAGCTCTGTGAA
TATCTGCTGTTCAAAGCTGGTGAAGCGGCGCCTTCAGCAGGGAAAAGTGACAATTGCTGA
GCTT [ATCCAGCCTGCTACACGGAGCCAG] GTGGATGCCACAGAACCACAGGATGTAGTGGT
GTATGACCAGAGCACACGAGATGCCAGCGTGCTGGCAGCAGACAGCTTCCTGTCCATCCT
GCTCAGCAAGCTGGACGGCTGCTTCGACAGTGTGGCCATCCTCACAGGAGGCTTCGCCAC
CTTCTCCTCCTGCTTCCCTGGCCTCTGTGAGGGCAAGCCTGCCACTCTACCGTCCATGAG
CCTCTCTCAGCCCTGCCTGCCTGTGCCAGTGTGGCCTGACCCGAATCCTGCCTCACCT
CTACCTGGGCTCTCAGAAAGATGTCTTGAACAAGGATCTGATGACCCAAAACGGAATAAG
CTATGTCTCAATGCCAGCAACTCCTGCCCTAAACCGGACTTCATCTGTGAGAGCCGTTT
CATGCGTATCCCCATCAATGACAATACTGTGAAAAGCTGCTGCCCTGGCTGGACAAGTC
CATCGAGTTTATTGATAAAGCCAAGCTGTCCAGCTGCCAAGTCATTGTTCACTGTCTGGC
TGGCATCTCTCGCTCTGCCACCATTGCCATCGCGTACATCATGAAAACCATGGGCATGTC
TTCTGACGACGCATACAGGTTTGTGAAGGATCGGCGCCCCCTCCATCTCGCCCCAACTTCAA
CTTCCTGGGCCAGTTGCTGGAGTATGAGAGGAGTCTGAAGCTGCTGGCTGCCCTGCAGAC
TGATGGACCTCACTTGGGGACCCCTGAGCCCCCTCATGGGCCCCGGCAGCAGGCATCCCACT
GCCCCGGCTGCCACCATCTACCTCAGAGAGCGCTGCCACTGGGAGCGAGGCAGCCACCGC
AGCCAGGGAGGGCAGCCCCAAGTGCTGGAGGGGATGCTCCGATCCCCAGCACAGCTCCAGC
CACCAGCGCGCTGCAGCAGGGCCTGCGTGCCCTGCACCTCTCCTCTGACCGCCTCCAGGA
CACCAACCGCCTCAAGCGTTTCTTTTCCCTGGACATCAAGTCGGCCTATGCACCCAGCAG
GAGGCCCGACTTTCCCGGCCACCCGACCCCGGTGAAGCCCCGAAGCTCTGCAAGCTGGA
CAGCCCGTCTGGGGGCACACTGGGCCTGCCCTCGCCCAGCCCAGACAGCCCGGACTCCGT
TCCAGAGTGCCGCCCACGACCCCGCCGGCGACGCCCCCGGCTAGTTTCGCCTGCCCGCTC
CCCCGCGCATGGTCTGGGCCTGAACTTTGGAGACACGGCCCCGGCAGACTCCACGGCACGG
CCTCTCGGCCCTGTGCGCGCCCGGGCTGCCCTGGCCCTGGCCAGCCGGCTGGCCCCGGGGG
CTGGGTGCCGCCACTGGACTCCCCAGGCACACCGTCGCCCGACGGCCCCCTGGTGCTTCAG
CCCCGAGGGCGCGCAGGGTCCAGGCGCTGTGTTCTCCGCCTTTGGCCGGGTAAGTGCAGG
CGCACCTGGACCCGGTAACAGCAGCAGCAGCGGTGGTGGTGGTGGTGGTGGTGGCGGCGG
CGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGG
CAGCAGCAGCAGCAGCAGCAGTAGTAGTAGTAGTAGTGACCTGCGGAGGCGGGATGTGCG
GACCGGCTGGCCCCGAGGAGCCTGCTGCAGATGCACAGTTCAAGAGGCGCAGCTGCCAGAT
GGAGTTCGAAGAGGGCATGGTGGAGGGGCGGGCACGTGGCGAGGAGCTGGCAGCCCTGGG
CAAGCAAACCAGCTTCTCTGGCAGCGTGGAGGTCATCGAAGTATCGTGACCCTTCAGAAG
TCCCTGTGCCCTTGCTCCAGCCAGGCCAGGTATAAATATATATTATATATAAAACACACA
GAAAAGGTAAATGGTTTTACTGCAATTTTTATCAAGAAGTAAATATTTTCGATTTTTTATT
TATTTAAGCTAGTGATCTGGCAACTGTGCGGGGCGGCCCTAAAGCTCTGTTTTTACTGTC
TGGTATTTAAACTGAAACAGGTTTCTAAGCAATATGAGGCCACCTTCAATCCCAAACCTGG
GTTGACAGGCCTGGGCCCCCTCCTTGCCCCCTCCCCTCTGGAAACATTACTGACCTTTCAA
GAGCTGCCCAGCTTTCCTGCACTTTTACATAAGAAAAAAGGGGGGGGGGGAA

FIGURE 2A

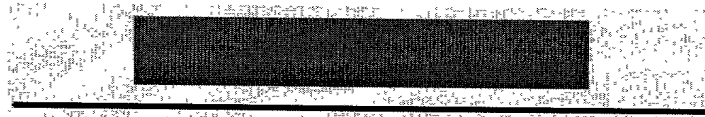
Gene Sequence Structure *

143 bp

Sequence Deleted

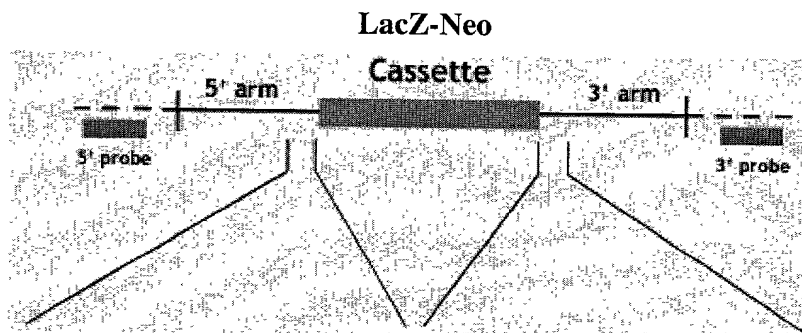
304 bp

Size of full-length cDNA: 2453 bp



Targeting Vector* (genomic sequence)

Arm Length:
5': 2.5 kb
3': 2 kb



Targeting Vector
- - - Endogenous Locus
* Not drawn to scale

<p>5' >TCCTGGGAGCCAGCTATAGCT ACCCAGATCCCACCATCTGCTGAC TATTCACCTTTCCCCCAGGTCCTGG CACCATGCACTAGGATACCCAGAA CGCTGCAAGGCCACGCCCTCCTCA CTTCAGGGGTCACCTCCTCCCATG CCCACCACCCACCATGGCTGGGG ATCGGCTCCCAGGAAGGTGATGG ACGCAAAGAAA<3' (SEQ ID NO:2)</p>	<p>5' >ATCCAGCCTGCTACACGAAGC CAGGTACCTGTGGCCCCACCCTTGC ATGCGTCTTCAGGGCTGACCATTG CTGAGCAAACAGACCTATGTCACC TCTGAAAGAGACAGAGGAGCTCCC AGGCCTGGTGCCAAGAGTCCTCTG ATAAGGCATTTCCCCCTCGCTGTC CCTCCGTTCCAAACAGGGTTCCTT GGGGTCAGAGC<3' (SEQ ID NO:3)</p>
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FIGURE 2B